0230







OTPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/007,343

DATE: 05/30/2002

TIME: 08:40:53

Input Set : A:\19202DB.txt

Output Set: N:\CRF3\05302002\J007343.raw

```
4 <110> APPLICANT: Linemeyer, David L.
              Hess, John W.
              Borkowski, Joseph A.
      6
              Bierilo, Kathleen K.
              Menke, John G.
     8
     10 <120> TITLE OF INVENTION: DNA ENCODING BRADYKININ B1 RECEPTOR
     13 <130> FILE REFERENCE: 19202DB
     15 <140> CURRENT APPLICATION NUMBER: 10/007,343
C--> 16 <141> CURRENT FILING DATE: 2002-05-17
     18 <150> PRIOR APPLICATION NUMBER: 09/307,912
     19 <151> PRIOR FILING DATE: 1999-05-10
     21 <150> PRIOR APPLICATION NUMBER: 08/932,761
     22 <151> PRIOR FILING DATE: 1997-09-17
     24 <150> PRIOR APPLICATION NUMBER: 08/229,418
     25 <151> PRIOR FILING DATE: 1994-04-15
     27 <160> NUMBER OF SEQ ID NOS: 2
     29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     31 <210> SEQ ID NO: 1
     32 <211> LENGTH: 1307
     33 <212> TYPE: DNA
     34 <213> ORGANISM: Human
     36 <400> SEQUENCE: 1
     37 cagagaaaac tcctccaaaa gcagctctca ctatcagaaa acccaactac agttgtgaac
                                                                                 60
     38 gccttcattt tctgcctgag gtctcagtcc gtcggcccag actgaagtgc agtggcacaa
                                                                               120
     39 tcatagctcg ctgcagcctc gaccttccag gcttaaacga ttctcccacc tcagcctctc
                                                                               180
                                                                               240
     40 gagttgctgg gaccacaggt cactgtgcat ggcatcatcc tggccccctc tagagctcca
     41 atcctccaac cagagccage tettecetca aaatgctaeg geetgtgaca atgctccaga
                                                                               300
     42 agcctgggac ctgctgcaca gagtgctgcc gacatttatc atctccatct gtttcttcgg
                                                                               360
     43 cctcctaggg aacctttttg tcctgttggt cttcctcctg ccccggcggc aactgaacgt
                                                                               420
     44 ggcagaaatc tacctggcca acctggcagc ctctgatctg gtgtttgtct tgggcttgcc
                                                                               480
     45 cttctgggca gagaatatct ggaaccagtt taactggcct ttcggagccc tcctctgccg
                                                                               540
     46 tgtcatcaac ggggtcatca aggccaattt gttcatcagc atcttcctgg tggtggccat
                                                                               600
     47 cagccaggac cgctaccgcg tgctggtgca ccctatggcc agcggaaggc agcagcggcg
                                                                                660
     48 gaggcaggcc cgggtcacct gcgtgctcat ctgggttgtg gggggcctct tgagcatccc
                                                                                720
     49 cacatteetg etgegateea tecaageegt eccagatetg aacateaceg eetgeateet
                                                                               780
     50 gctcctccc catgaggcct ggcactttgc aaggattgtg gagttaaata ttctgggttt
                                                                               840
     51 cctcctacca ctggctgcga tcgtcttctt caactaccac atcctggcct ccctgcgaac
                                                                               900
                                                                               960
     52 gegggaggag gteageagga caaggtgegg gggeegeaag gatageaaga eeacageget
     53 gatecteacg etegtggttg cetteetggt etgetgggee cettaceact tetttgeett
                                                                              1020
     54 cctggaattc ttattccagg tgcaagcagt ccgaggctgc ttttgggagg acttcattga
                                                                              1080
     55 cctgggcctg caattggcca acttctttgc cttcactaac agctccctga atccagtaat
                                                                               1140
                                                                              1200
     56 ttatgtcttt gtgggccggc tcttcaggac caaggtctgg gaactttata aacaatgcac
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     57 ccctaaaagt cttgctccaa tatcttcatc ccataggaaa gaaatcttcc aacttttctg
```

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58	acad	aatt	caa a	acac	catt	g aa	ccaa	gaaa	a aaa	aaaa	aaa	aaaa	ıaaa				1307
		•		NO:													
61	<213	L> LI	ENGTE	I: 35	3												
62	<212	2> T	PE:	PRT													
63	<213	3> OI	RGANI	SM:	Huma	an											
65	<400)> SI	EQUE	ICE:	2 ·												
66	Met	Ala	Ser	Ser	${\tt Trp}$	Pro	Pro	Leu	Glu	Leu	Gln	Ser	Ser	Asn	Gln	Ser	
67	1				5					10					15		
68	Gln	Leu	Phe	Pro	Gln	Asn	Ala	Thr	Ala	Cys	Asp	Asn	Ala	Pro	Glu	Ala	
69				20					25					30			
70	Trp	Asp	Leu	Leu	His	Arg	Val	Leu	Pro	Thr	Phe	Ile	Ile	Ser	Ile	Cys	
71			35					40					45				
72	Phe	Phe	Gly	Leu	Leu	Gly	Asn	Leu	Phe	Val	Leu	Leu	Val	Phe	Leu	Leu	
73		50					55 _.					60					
74	Pro	Arg	Arg	Gln	Leu	Asn	Val	Ala	Glu	Ile	Tyr	Leu	Ala	Asn	Leu	Ala	
75	65					70					75					80	
76	Ala	Ser	Asp	Leu	Val	Phe	Val	Leu	Gly	Leu	Pro	Phe	${\tt Trp}$	Ala	Glu	Asn	
77					85					90					95		
78	Ile	Trp	Asn	Gln	Phe	Asn	Trp	Pro	Phe	Gly	Ala	Leu	Leu	Cys	Arg	Val	
79				100					105					110			
80	Ile	Asn	Gly	Val	Ile	Lys	Ala	Asn	Leu	Phe	Ile	Ser	Ile	Phe	Leu	Val	
81			115					120					125				
82	Val	Ala	Ile	Ser	Gln	Asp	Arg	Tyr	Arg	Val	Leu	Val	His	Pro	Met	Ala	
83		130					135					140					
84	Ser	Gly	Arg	Gln	Gln	Arg	Arg	Arg	Gln	Ala	Arg	Val	Thr	Cys	Val	Leu	
85	145					150					155					160	
86	Ile	Trp	Val	Val	Gly	Gly	Leu	Leu	Ser	Ile	Pro	Thr	Phe	Leu	Leu	Arg	
87					165					170					175		
	Ser	Ile	Gln		Val	Pro	Asp	Leu		Ile	Thr	Ala	Cys	Ile	Leu	Leu	
89				180					185					190			
	Leu	Pro		Glu	Ala	Trp	His		Ala	Arg	Ile	Val		Leu	Asn	Ile	
91		_	195					200		_	0_		205				
	Leu		Phe	Leu	Leu	Pro		Ala	Ala	Ile	Val		Phe	Asn	\mathtt{Tyr}	His	
93		210					215					220					
		Leu	Ala	Ser	Leu		Thr	Arg	GLu	GLu		Ser	Arg	Thr	Arg		
	225				_	230					235		_		_	240	
	GLY	Gly	Arg	Lys	_	Ser	Lys	Thr	Thr		Leu	Ile	Leu	Thr	Leu	Val	
97			_,	_	245	_	_		_	250	•	1	1	- 1	255	_	
															Phe	Leu	
											a 1					_	
		ı Phe			: GIr	ı vaı	. GIn			. Arg	l GTÀ	Cys			Glu	Asp	
101		-1	275		~ 1	.	a1 .	280			n1	D1 -	285		ml.		
			_) ren	GIY	Leu					Pne			Pne	Thr	ASn	
103		290		7	. D	77-7	295		. 17. 1		17-7	300		 .	. DL -	3	
			. Leu	ASI	PIC			туг	val	. Pue			AIC	, тег	ı Phe		
	305		. 37-7	П		310		T		. a	315		т		. T	320	
		. гуд	s val	. Trp	325		туг	пλε	GIN	330		PIO	гλε	sei	Leu 335		
107		. ті-					A	т	. dl			. cı -	т	nh-			
TAR	PLC	TTE	: ser	ser	sei	nıs	AI.d	гÀг	GIU	тте	: Flie	: GII	ьес	r bile	Trp	Arg	

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345

109 110 Asn VERIFICATION SUMMARY

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L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date